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RAW SEQUENCE LISTING

DATE: 05/16/2002

PATENT APPLICATION: US/10/024,368

TIME: 14:13:08

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1 <110> APPLICANT: THOMPSON, Catherine C.
 2 <120> TITLE OF INVENTION: HUMAN HAIRLESS GENE AND PROTEIN
 3 <130> FILE REFERENCE: Thompson-20263/0243435
 4 <140> CURRENT APPLICATION NUMBER: 10/024,368
 5 <141> CURRENT FILING DATE: 2001-12-21
 7 <150> PRIOR APPLICATION NUMBER: EARLIER APPLICATION NUMBER: US/09/287,354
 W--> 8 <151> PRIOR FILING DATE: EARLIER FILING DATE: 1999-04-07
 11 <150> PRIOR APPLICATION NUMBER: EARLIER APPLICATION NUMBER: US 60/080,888
 W--> 12 <151> PRIOR FILING DATE: EARLIER FILING DATE: 1998-04-07
 13 <160> NUMBER OF SEQ ID NOS: 10
 14 <170> SOFTWARE: PatentIn Ver. 2.0
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 17 <211> LENGTH: 3202
 18 <212> TYPE: DNA
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77 <211> LENGTH: 984

78 <212> TYPE: PRT

79 <213> ORGANISM: Homo sapiens

80 <400> SEQUENCE: 2

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84 20 25 30
85 Gln Arg Ala Gly Glu Ala Glu Arg Pro Ser Leu His Gln Arg Asp Gly
86 35 40 45
87 Glu Met Gly Ala Gly Arg Gln Asn Pro Cys Pro Leu Phe Leu Gly
88 50 55 60
89 Gln Pro Asp Thr Val Pro Trp Thr Ser Trp Pro Ala Cys Pro Pro Gly
90 65 70 75 80
91 Leu Val His Thr Leu Gly Asn Val Trp Ala Gly Pro Gly Asp Gly Asn
92 85 90 95
93 Leu Gly Tyr Gln Leu Gly Pro Pro Ala Thr Pro Arg Cys Pro Ser Pro
94 100 105 110
95 Glu Pro Pro Val Thr Gln Arg Gly Cys Cys Ser Ser Tyr Pro Pro Thr
96 115 120 125
97 Lys Gly Gly Asp Leu Gly Pro Cys Gly Lys Cys Gln Glu Gly Leu Glu

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101  Ser Gly Pro Arg Ala Cys Pro Pro Ser His His Thr Lys Leu Lys Lys
102      165      170      175
103  Thr Trp Leu Thr Arg His Ser Glu Gln Phe Glu Cys Pro Arg Gly Cys
104      180      185      190
105  Pro Glu Val Glu Glu Arg Pro Val Ala Arg Leu Arg Ala Leu Lys Arg
106      195      200      205
107  Ala Gly Ser Pro Glu Val Gln Gly Ala Met Gly Ser Pro Ala Pro Lys
108      210      215      220
109  Arg Pro Pro Asp Pro Phe Pro Gly Thr Ala Glu Gln Gly Ala Gly Gly
110      225      230      235      240
111  Trp Gln Glu Val Arg Asp Thr Ser Ile Gly Asn Lys Asp Val Asp Ser
112      245      250      255
113  Gly Gln His Asp Glu Gln Lys Gly Pro Gln Asp Gly Gln Ala Ser Leu
114      260      265      270
115  Gln Asp Pro Gly Leu Gln Asp Ile Pro Cys Leu Ala Leu Pro Ala Lys
116      275      280      285
117  Leu Ala Gln Cys Gln Ser Cys Ala Gln Ala Ala Gly Glu Gly Gly Gly
118      290      295      300
119  His Ala Cys His Ser Gln Val Arg Arg Ser Pro Leu Gly Gly Glu
120      305      310      315      320
121  Leu Gln Gln Glu Glu Asp Thr Ala Thr Asn Ser Ser Ser Glu Glu Gly
122      325      330      335
123  Pro Gly Ser Gly Pro Asp Ser Arg Leu Ser Thr Gly Leu Ala Lys His
124      340      345      350
125  Leu Leu Ser Gly Leu Gly Asp Arg Leu Cys Arg Leu Leu Arg Arg Glu
126      355      360      365
127  Arg Glu Ala Leu Ala Trp Ala Gln Arg Glu Gly Gln Gly Pro Ala Val
128      370      375      380
129  Thr Gly Asp Ser Pro Gly Ile Pro Arg Cys Cys Ser Arg Cys His His
130      385      390      395      400
131  Gly Leu Phe Asn Thr His Trp Arg Cys Pro Arg Cys Ser His Arg Leu
132      405      410      415
133  Cys Val Ala Cys Gly Arg Val Ala Gly Thr Gly Arg Ala Arg Glu Lys
134      420      425      430
135  Ala Gly Phe Gln Glu Gln Ser Ala Glu Glu Cys Thr Gln Glu Ala Gly
136      435      440      445
137  His Ala Ala Cys Ser Leu Met Leu Thr Gln Phe Val Ser Ser Gln Ala
138      450      455      460
139  Leu Ala Glu Leu Ser Thr Ala Met His Gln Val Trp Val Lys Phe Asp
140      465      470      475      480
141  Ile Arg Gly His Cys Pro Cys Gln Ala Asp Ala Arg Val Trp Ala Pro
142      485      490      495
143  Gly Asp Ala Gly Gln Gln Lys Glu Ser Thr Gln Lys Thr Pro Pro Thr
144      500      505      510
145  Pro Gln Pro Ser Cys Asn Gly Asp Thr His Arg Thr Lys Ser Ile Lys
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150	545					550					555					560
151	Ala	Val	Lys	Leu	Cys	Leu	Gly	His	Glu	Arg	Ile	His	Met	Ala	Phe	Ala
152					565					570						575
153	Pro	Val	Thr	Pro	Ala	Leu	Pro	Ser	Asp	Arg	Ile	Thr	Asn	Ile	Leu	
154				580					585					590		
155	Asp	Ser	Ile	Ala	Gln	Val	Val	Glu	Arg	Lys	Ile	Gln	Glu	Lys	Ala	
156			595					600					605			
157	Leu	Gly	Pro	Gly	Leu	Arg	Ala	Gly	Pro	Gly	Leu	Arg	Lys	Gly	Leu	Gly
158		610					615					620				
159	Leu	Pro	Leu	Ser	Pro	Val	Arg	Pro	Arg	Leu	Pro	Pro	Pro	Gly	Ala	Leu
160	625					630					635					640
161	Leu	Trp	Leu	Gln	Glu	Pro	Gln	Pro	Cys	Pro	Arg	Arg	Gly	Phe	His	Leu
162					645						650					655
163	Phe	Gln	Glu	His	Trp	Arg	Gln	Gly	Gln	Pro	Val	Leu	Val	Ser	Gly	Ile
164				660					665						670	
165	Gln	Arg	Thr	Leu	Gln	Gly	Asn	Leu	Trp	Gly	Thr	Glu	Ala	Leu	Gly	Ala
166			675					680					685			
167	Leu	Gly	Gly	Gln	Val	Gln	Ala	Leu	Ser	Pro	Leu	Gly	Pro	Pro	Gln	Pro
168		690					695					700				
169	Ser	Ser	Leu	Gly	Ser	Thr	Phe	Trp	Glu	Gly	Phe	Ser	Trp	Pro	Glu	
170	705					710					715					720
171	Leu	Arg	Pro	Lys	Ser	Asp	Glu	Gly	Ser	Val	Leu	Leu	Leu	His	Arg	Ala
172					725					730						735
173	Leu	Gly	Asp	Glu	Asp	Thr	Ser	Arg	Val	Glu	Asn	Leu	Ala	Ala	Ser	Leu
174				740					745					750		
175	Pro	Leu	Pro	Glu	Tyr	Cys	Ala	Leu	His	Gly	Lys	Leu	Asn	Leu	Ala	Ser
176			755					760					765			
177	Tyr	Leu	Pro	Pro	Gly	Leu	Ala	Leu	Arg	Pro	Leu	Glu	Pro	Gln	Leu	Trp
178		770					775					780				
179	Ala	Ala	Tyr	Gly	Val	Ser	Pro	His	Arg	Gly	His	Leu	Gly	Thr	Lys	Asn
180	785					790					795					800
181	Leu	Cys	Val	Glu	Val	Ala	Asp	Leu	Val	Ser	Ile	Leu	Val	His	Ala	Asp
182					805					810						815
183	Thr	Pro	Leu	Pro	Ala	Trp	His	Arg	Ala	Gln	Lys	Asp	Phe	Leu	Ser	Gly
184				820					825						830	
185	Leu	Asp	Gly	Glu	Gly	Leu	Trp	Ser	Pro	Gly	Ser	Gln	Val	Ser	Thr	Val
186			835					840					845			
187	Trp	His	Val	Phe	Arg	Ala	Gln	Asp	Ala	Gln	Arg	Ile	Arg	Arg	Phe	Leu
188		850					855						860			
189	Gln	Met	Val	Cys	Pro	Ala	Gly	Ala	Gly	Ala	Leu	Glu	Pro	Gly	Ala	Pro
190	865					870					875					880
191	Gly	Ser	Cys	Tyr	Leu	Asp	Ala	Gly	Leu	Arg	Arg	Arg	Leu	Arg	Glu	Glu
192					885					890					895	
193	Trp	Gly	Val	Ser	Cys	Trp	Thr	Leu	Leu	Gln	Ala	Pro	Gly	Glu	Ala	Val
194				900						905					910	
195	Leu	Val	Pro	Ala	Gly	Ala	Pro	His	Gln	Val	Gln	Gly	Leu	Val	Ser	Thr

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198          930          935          940
199 Ala Gln Leu Cys His Gln Gly Pro Ser Leu Pro Pro Asp Cys His Leu
200          945          950          955          960
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204          980
206 <210> SEQ ID NO: 3
207 <211> LENGTH: 1189
208 <212> TYPE: PRT
209 <213> ORGANISM: Homo sapiens
210 <400> SEQUENCE: 3
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215 Arg Asp Gly Leu His His Gly Pro Leu Cys Leu Gly Glu Pro Ala Pro
216          35          40          45
217 Phe Trp Arg Gly Val Leu Ser Thr Pro Asp Ser Trp Leu Pro Pro Gly
218          50          55          60
219 Phe Pro Gln Gly Pro Lys Asp Met Leu Pro Leu Val Glu Gly Glu Gly
220          65          70          75          80
221 Pro Gln Asn Gly Glu Arg Lys Val Asn Trp Leu Gly Ser Lys Glu Gly
222          85          90          95
223 Leu Arg Trp Lys Glu Ala Met Leu Thr His Pro Leu Ala Phe Cys Gly
224          100          105          110
225 Pro Ala Cys Pro Pro Arg Cys Gly Pro Leu Met Pro Glu His Ser Gly
226          115          120          125
227 Gly His Leu Lys Ser Asp Pro Val Ala Phe Arg Pro Trp His Cys Pro
228          130          135          140
229 Phe Leu Leu Glu Thr Lys Ile Leu Glu Arg Ala Pro Phe Trp Val Pro
230          145          150          155          160
231 Thr Cys Leu Pro Pro Tyr Leu Val Ser Gly Leu Pro Pro Glu His Pro
232          165          170          175
233 Cys Asp Trp Pro Leu Thr Pro His Pro Trp Val Tyr Ser Gly Gly Gln
234          180          185          190
235 Pro Lys Val Pro Ser Ala Phe Ser Leu Gly Ser Lys Gly Phe Tyr Tyr
236          195          200          205
237 Lys Asp Pro Ser Ile Pro Arg Leu Ala Lys Glu Pro Leu Ala Ala Ala
238          210          215          220
239 Glu Pro Gly Leu Phe Gly Leu Asn Ser Gly Gly His Leu Gln Arg Ala
240          225          230          235          240
241 Gly Glu Ala Glu Arg Pro Ser Leu His Gln Arg Asp Gly Glu Met Gly
242          245          250          255
243 Ala Gly Arg Gln Gln Asn Pro Cys Pro Leu Phe Leu Gly Gln Pro Asp
244          260          265          270
245 Thr Val Pro Trp Thr Ser Trp Pro Ala Cys Pro Pro Gly Leu Val His

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VERIFICATION SUMMARY

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